

Grid Computing to Study the Functions of Plant Genes

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A Bit of Background

- A major goal in biology is to learn the function of each gene in an organism.
- A proven approach is to compare the behaviors of individuals possessing different versions of that gene.
- Organisms have on the order of 10⁴ genes so that makes for a lot of comparisons.



25,000 genes

Α 0.40 0.35 cry1 Growth Rate (πm min -1 0.20 0.20 0.15 0.10 0.05 0.00 0 2 3 5 6 Λ Time (h)

WT

cry1

Hypocotyl Growth Inhibition Induced by Blue Light

We developed image processing algorithms for measuring seedling growth and development in order to quantify effects of genetic differences (i.e. phenotypes of mutants) in space and time. Our purpose was to answer questions about one or two genes.

Why not / How to scale up to the whole genome level?



Alonso-Blanco et al. The Plant Journal 14, 259-257 (1998)

Let's make the process High Throughput Switch to Root Gravitropism...

because we could study with high resolution, high accuracy, and high throughput



Machine Vision to Study Natural Genetic Variation

160 Ler X CVI recombinant inbred lines for QTL mapping



A single QTL plot (for example - root tip angle at one point in time is the phenotype)







The Key People



Logan Johnson

Candace Moore

Corn is king









Huge diversity in ear shapes among genetically-distinct maize types



Is this the best way to measure corn ear 'traits'?



Too funny –

"Geneticist <u>Ed Buckler</u> measures a maize ear for statistical analysis."

Corn ear and kernel traits measured by image analysis



The current corn ear and kernel workflow





Image stacks

Biology is becoming a high-throughput science

New insights will come from Big Data

People who are comfortable with HTC will win!